

10/520881

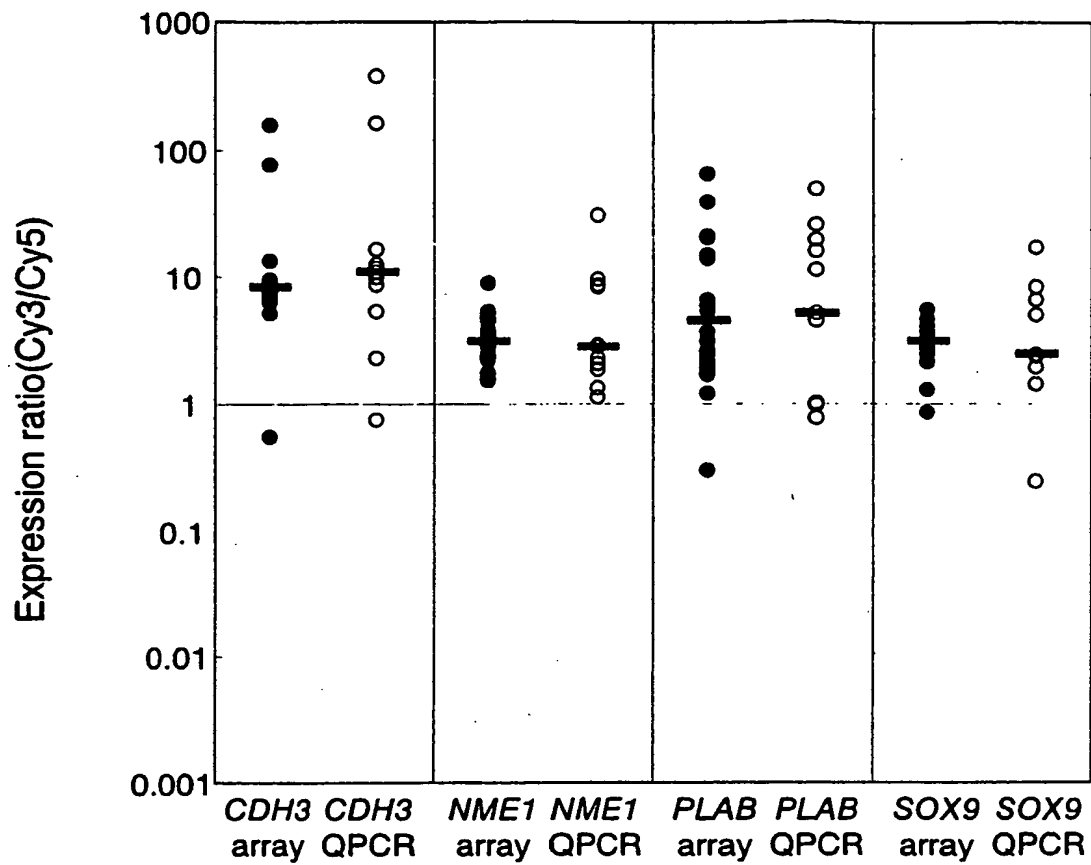


Fig. 1

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Fig. 2A

Symbol	Title	GenBank Accession	training												test		permutational P	discriminant coefficient																		
			node-positive						node-negative						node positive	node negative																				
			104+	75+	72+	63+	61+	59+	58+	51+	48+	47+	57+	73+	97+	101+	102+	134+	135+	136+	137+	138+														
DDOST	dolichyl-diphosphooligosaccharide- protein glycosyltransferase	BF941738	0.83	1.63	-0.23	-0.07	1.36	0.41	0.72	1.75	0.36	0.02	-0.06	-0.13	2.0	-0.26	0.86	0.72	-1.08	1.28	-0.39	0.44	0.25	0.63	0.59	-0.49				-0.93				0.00006	1.87	
GNS	glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	Z12173	1.42	2.56	-0.10	0.09	1.50	-0.14	0.81	1.59	1.09	-0.14	0.18	-0.11	3.3	0.86	0.72	-1.08	1.28	-0.39	0.44	0.25	1.04	0.67	-0.35				-0.39				0.00119	1.26		
NEDD8	neural precursor cell expressed, developmentally down-regulated 8	D23662	2.23	0.70	0.23	0.59	3.65	-0.09	0.60	0.24	0.53	-1.25	-0.49	-0.36	4.0	1.03	0.07	-0.67	-0.34	1.25	-0.24	0.13	-0.09	0.24	0.35	1.11	0.71		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
	DKFZP5661133	AA533633	1.32	1.28	0.29	0.75	1.27	0.69	1.60	0.68	1.15	0.29	-0.20	0.25	3.8	-0.26	0.86	0.72	-1.08	1.28	-0.39	0.44	0.25	0.20	0.53	-0.56	-0.14	-0.48	0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
	ESTs, Highly similar to B27079 fibronectin receptor beta chain precursor	A1755112	1.20	0.56	-0.14	1.33	2.91	0.95	0.60	0.43	-0.02	-0.29	-0.45	-0.51	2.8	0.49	1.31	0.01	-0.26	1.43	-0.39	0.74	1.00	0.51	0.37	0.84	0.72		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
LOC 51096	CGI-48 protein	AA448503	0.82	1.76	1.18	-0.39	1.31	0.13	2.03	1.24	1.60	-0.61	-0.77	-1.42	4.7	0.20	0.53	-0.56	-0.14	-0.48	0.09	0.70	0.12	-0.45	0.21	0.85	1.11		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
CCT5	chaperonin containing TCP1, subunit 5 (epsilon)	D43950	1.60	0.97	1.16	0.80	2.37	-0.51	1.52	0.25	0.73	-0.13	0.10	-0.43	2.9	0.49	1.31	0.01	-0.26	1.43	-0.39	0.74	1.00	0.51	0.37	0.84	0.72		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
CCT3	chaperonin containing TCP1, subunit 3 (gamma)	X74801	0.68	1.93	1.01	0.97	1.26	0.04	2.04	1.14	1.25	-0.73	0.81	-1.00	1.9	0.20	0.53	-0.56	-0.14	-0.48	0.09	0.70	0.12	-0.45	0.21	0.85	1.11		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
PPP 2R1B	protein phosphatase 2 (formerly 2A), regulatory subunit A (PR65), beta isoform	M65254	1.51	1.29	0.19	1.79	2.94	-0.66	0.57	0.50	1.01	-0.34	-1.09	-0.35	3.5	0.20	0.53	-0.56	-0.14	-0.48	0.09	0.70	0.12	-0.45	0.21	0.85	1.11		0.09	0.70	0.12	-0.45	0.21	0.85	1.11	-2.8
UBQLN1	ubiquitin 1	AA281115																																		
AIM2	absent in melanoma 2	AF024714																																		
USP9X	ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)	X98296																																		
			"Predictive score"																																	
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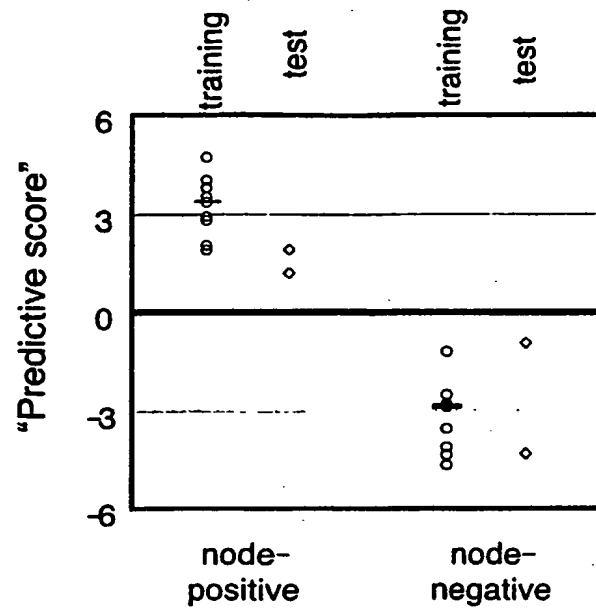


Fig. 2B

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